

SEQUENCE LISTING

<110> Patten, Phillip A. et al.

<120> Interferon-Alpha Polypeptides and Conjugates

<130> 0269us410

<150> US 10/714,817

<151> 2003-11-17

<150> US 60/502,560

<151> 2003-09-12

<150> US 60/427,612

<151> 2002-11-18

<160> 104

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<212> PRT

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<220>

<223> IFNalpha B9x11

<400> 1

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
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Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35				40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55					60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115				120						125			
Leu	Tyr	Leu	Thr	Lys	Lys	Lys	Tyr	Ser	Pro	Cys	Ser	Trp	Glu	Val	Val
	130				135						140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
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<213> Artificial Sequence

<220>

<223> IFNalpha B9x12

<400> 2

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
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Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Lys	Lys	Lys	Tyr	Ser	Pro	Cys	Ser	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
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<220>

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<400> 3

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
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Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
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<210> 4
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 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Phe His Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Lys Lys Lys Tyr Ser Pro Cys Ser Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys
 145 150 155 160
 Arg Leu Arg Arg Lys Glu
 165

<210> 5
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 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x16

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 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val

130		135		140											
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145		150		155											160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 6
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<220>
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Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Phe	His	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
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<210> 7
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 <223> IFNalpha B9x18

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Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe
		35					40					45			
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Phe	His	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	

Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 8
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x21

<400>	8														
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1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25				30			
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe
		35				40					45				
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr
	50					55				60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
				85					90				95		
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 9
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 <212> PRT
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<220>
 <223> IFNalpha B9x22

<400>	9														
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25				30			
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35				40					45				
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr

50		55		60	
Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr					
65		70		75	80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu					
	85		90		95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met					
	100		105		110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr					
	115		120		125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val					
	130		135		140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys					
145		150		155	160
Arg Leu Arg Arg Lys Glu					
	165				

<210> 10
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x23

<400> 10	
Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met	
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Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp	
	20 25 30
Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe	
	35 40 45
Gln Lys Val Gln Ala Ile Phe Leu Leu Tyr Glu Leu Ile Gln Gln Thr	
	50 55 60
Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr	
65	70 75 80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu	
	85 90 95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met	
	100 105 110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr	
	115 120 125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val	
	130 135 140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys	
145	150 155 160
Arg Leu Arg Arg Lys Glu	
	165

<210> 11
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x24

<400> 11	
Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met	
1	5 10 15

Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
		20						25				30			
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe
		35					40					45			
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Lys	Lys	Lys	Tyr	Ser	Pro	Cys	Ser	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 12
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5				10					15		
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
		20						25				30			
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 13
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 <212> PRT
 <213> Artificial Sequence

<220>

<223> IFNalpha B9x26

<400> 13

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
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Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35					40					45			
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Lys	Lys	Lys	Tyr	Ser	Pro	Cys	Ser	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
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<210> 14

<211> 166

<212> PRT

<213> Artificial Sequence

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<223> IFNalpha B9x27

<400> 14

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Lys	Lys	Lys	Tyr	Ser	Pro	Cys	Ser	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
					165										

<210> 15

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<220>
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 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Leu Tyr Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Lys Lys Lys Tyr Ser Pro Cys Ser Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys
 145 150 155 160
 Arg Leu Arg Arg Lys Glu
 165

<210> 16
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<220>
 <223> IFNalpha B9x11 coding sequence

<400> 16
 tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacaa 60
 atgaggagaa tctctctttt ctcctgtctg aaggacagac atgacttcag atttccccag 120
 gaggagtgtg atggcaacca cttccagaag gttcaagcta tcttcctttt ctatgagatg 180
 atgcagcaga ccttcaacct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
 ctccatagaaa aattctacat tgaacttttc cagcaaataga atgacctgga agcctgcgtg 300
 atgcaggagg ttggagtggg agagactccc ctgatgaatg tggactccat cctggctgtg 360
 aggaaatact ttcaaagaat cactctttat ctgacaaaaga agaagtatag cccttggtcc 420
 tgggagggtg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
 agattaagga ggaaggaa 498

<210> 17
 <211> 498
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x12 coding sequence

<400> 17
 tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacaa 60

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atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca gttccagaag gttcaagcta tcttcctttt ctatgagatg 180
atgcagcaga ccttcaacct ctccagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata atgacctgga agcctgcgtg 300
atgcaggagg ttggagtggg agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacaaaaga agaagtatag cccttggttc 420
tgggagggtt tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa
498

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<210> 18

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x14 coding sequence

<400> 18

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tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacia 60
atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca gttccagaag gttcaagcta tcttcctttt ctatgagatg 180
atgcagcaga ccttcaacct ctccagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata atgacctgga agcctgcgtg 300
atgcaggagg ttggagtggg agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacagaga agaagtatag cccttggttc 420
tgggagggtt tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa
498

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<210> 19

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x15 coding sequence

<400> 19

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tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacia 60
atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca gttccagaag actcaagcta tctctgtctt ccatgagatg 180
atgcagcaga ccttcaacct ctccagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata atgacctgga agcctgcgtg 300
atgcaggagg ttggagtggg agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacaaaaga agaagtatag cccttggttc 420
tgggagggtt tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa
498

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<210> 20

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x16 coding sequence

<400> 20

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tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacia 60
atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca gttccagaag gttcaagcta tcttcctttt ctatgagatg 180
atgcagcaga ccttcaacct ctccagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata atgacctgga agcctgcgtg 300

```

```

atgcaggagg ttggagtgga agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacagaga agaagtatag cccttgtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa

```

<210> 21

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x17 coding sequence

<400> 21

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tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacia 60
atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca cttccagaag actcaagcta tctctgtctt ccatgagatg 180
atgcagcaga cttcaacct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga atgacctgga agcctgcgtg 300
atgcaggagg ttggagtgga agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacagaga agaagtatag cccttgtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa

```

<210> 22

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x18 coding sequence

<400> 22

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tgtgatctgc ctcagaccca cagcctgggt cacaggagga ccatgatgct cctggcacia 60
atgaggagaa tctctctttt ctctgtctg aaggacagac atgacttcag atttccccag 120
gaggagtttg atggcaacca gttccagaag actcaagcta tctctgtctt ccatgagatg 180
atgcagcaga cttcaacct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga atgacctgga agcctgcgtg 300
atgcaggagg ttggagtgga agagactccc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact ttcaaagaat cactctttat ctgacagaga agaagtatag cccttgtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa

```

<210> 23

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x21 coding sequence

<400> 23

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tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacia 60
atgaggagaa tctctctttt ctctgcctg aaggacagac atgatttcgg atttccccag 120
gaggagtttg atggccacca gttccagaag actcaagcca tctctgtcct ccatgagctg 180
atccagcaga cttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga ataacctgga agcatgtgtg 300
atacaggagg ttgggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacagaga agaaatacag cccttgtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa

```

<210> 24
<211> 498
<212> DNA
<213> Artificial Sequence

<220>
<223> IFNalpha B9x22 coding sequence

<400> 24
tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctccctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca cttccagaag actcaagcca tctctgtcct ccatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata gaacactgga agcatgtgtg 300
atacaggagg ttgggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacagaga agaaatacag cccttggtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa 498

<210> 25
<211> 498
<212> DNA
<213> Artificial Sequence

<220>
<223> IFNalpha B9x23 coding sequence

<400> 25
tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctccctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca gttccagaag gttcaagcca tcttccttct ctatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata gaacactgga agcatgtgtg 300
atacaggagg ttgggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacagaga agaaatacag cccttggtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa 498

<210> 26
<211> 498
<212> DNA
<213> Artificial Sequence

<220>
<223> IFNalpha B9x24 coding sequence

<400> 26
tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctccctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca gttccagaag actcaagcca tctctgtcct ccatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaata gaacactgga agcatgtgtg 300
atacaggagg ttgggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacaaa gaatacag cccttggtcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa 498

<210> 27
<211> 498
<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x25 coding sequence

<400> 27

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tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca cttccagaag gttcaagcca tcttccttct ctatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga ataacctgga agcatgtgtg 300
atacaggagg ttggggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacagaga agaaatacag cccttggtgcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa                                498
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<210> 28

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x26 coding sequence

<400> 28

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tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca cttccagaag actcaagcca tctctgtcct ccatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga ataacctgga agcatgtgtg 300
atacaggagg ttggggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacaaaaga agaaatacag cccttggttcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa                                498
```

<210> 29

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x27 coding sequence

<400> 29

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tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
atgaggagaa tctctccttt ctctgcctg aaggacagac atgatttcgg attccccgag 120
gaggagtttg atggccacca gttccagaag gttcaagcca tcttccttct ctatgagctg 180
atccagcaga ctttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
ctcctagaaa aattctacat tgaacttttc cagcaaatga ataacctgga agcatgtgtg 300
atacaggagg ttggggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
aggaaatact tccgaagaat cactctctat ctgacaaaaga agaaatacag cccttggttcc 420
tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
agattaagga ggaaggaa                                498
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<210> 30

<211> 498

<212> DNA

<213> Artificial Sequence

<220>

<223> IFNalpha B9x28 coding sequence

<400> 30
 tgtgatctgc ctcagaccca cagcctgagt aacaggagga ctctgatgct catggcacaa 60
 atgaggagaa tctctccttt ctctgcctg aaggacagac atgatttcgg attccccgag 120
 gaggagtttg atggccacca cttccagaag gttcaagcca tcttccttct ctatgagctg 180
 atccagcaga ccttcaatct cttcagcaca aagaactcat ctgctgcttg ggatgagacc 240
 ctctagaaa aattctacat tgaacttttc cagcaaatga ataacctgga agcatgtgtg 300
 atacaggagg ttgggggtgga agagattgcc ctgatgaatg tggactccat cctggctgtg 360
 aggaaatact tccgaagaat cactctctat ctgacaaaga agaaatacag cccttggttcc 420
 tgggaggttg tcagagcaga aatcatgaga tctttctctt tttcaacaaa cttgcaaaaa 480
 agattaagga ggaaggaa 498

<210> 31
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-1a

<400> 31
 Cys Asp Leu Pro Glu Thr His Ser Leu Asp Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Leu Ala Gln Met Ser Arg Ile Ser Pro Ser Ser Cys Leu Met Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Ala Pro Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Ile
 50 55 60
 Phe Asn Leu Phe Thr Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Asp
 65 70 75 80
 Leu Leu Asp Lys Phe Cys Thr Glu Leu Tyr Gln Gln Leu Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Glu Arg Val Gly Glu Thr Pro Leu Met
 100 105 110
 Asn Ala Asp Ser Ile Leu Ala Val Lys Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Leu Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Arg Leu Arg Arg Lys Glu
 165

<210> 32
 <211> 165
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-2b

<400> 32
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly Ser Arg Arg Thr Leu Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Gln Glu Glu Phe Gly Asn Gln Phe Gln
 35 40 45
 Lys Ala Glu Thr Ile Pro Val Leu His Glu Met Ile Gln Gln Ile Phe

50		55		60											
Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr	Leu
65					70					75					80
Leu	Asp	Lys	Phe	Tyr	Thr	Glu	Leu	Tyr	Gln	Gln	Leu	Asn	Asp	Leu	Glu
				85						90				95	
Ala	Cys	Val	Ile	Gln	Gly	Val	Gly	Val	Thr	Glu	Thr	Pro	Leu	Met	Lys
			100					105					110		
Glu	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr	Leu
		115					120					125			
Tyr	Leu	Lys	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val	Arg
		130				135					140				
Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu	Ser
145					150					155					160
Leu	Arg	Ser	Lys	Glu											
				165											

<210> 33
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-4b

<400> 33
Cys Asp Leu Pro Gln Thr His Ser Leu Gly Asn Arg Arg Ala Leu Ile
1 5 10 15
Leu Leu Ala Gln Met Gly Arg Ile Ser His Phe Ser Cys Leu Lys Asp
20 25 30
Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
35 40 45
Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln Thr
50 55 60
Phe Asn Leu Phe Ser Thr Glu Asp Ser Ser Ala Ala Trp Glu Gln Ser
65 70 75 80
Leu Leu Glu Lys Phe Ser Thr Glu Leu Tyr Gln Gln Leu Asn Asp Leu
85 90 95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
100 105 110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
115 120 125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
130 135 140
Arg Ala Glu Ile Met Arg Ser Leu Ser Phe Ser Thr Asn Leu Gln Lys
145 150 155 160
Arg Leu Arg Arg Lys Asp
165

<210> 34
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-5

<400> 34
Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
1 5 10 15

Ile	Met	Ala	Gln	Met	Gly	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe
		35					40					45			
Gln	Lys	Ala	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Thr	Trp	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Asp	Lys	Phe	Tyr	Thr	Glu	Leu	Tyr	Gln	Gln	Leu	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Met	Met	Gln	Glu	Val	Gly	Val	Glu	Asp	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Thr	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Ala	Asn	Leu	Gln	Glu
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 35
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-6

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25				30			
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe
		35					40					45			
Gln	Lys	Ala	Glu	Ala	Ile	Ser	Val	Leu	His	Glu	Val	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Val	Ala	Trp	Asp	Glu	Arg
65					70					75					80
Leu	Leu	Asp	Lys	Leu	Tyr	Thr	Glu	Leu	Tyr	Gln	Gln	Leu	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Trp	Val	Gly	Gly	Thr	Pro	Leu	Met
			100					105					110		
Asn	Glu	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Ser	Ser	Arg	Asn	Leu	Gln	Glu
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 36
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>

<223> mature huIFN alpha-7a

<400> 36

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Arg	Asn	Arg	Arg	Ala	Leu	Ile
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Gly	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Glu	Phe	Arg	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe
		35					40					45			
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Glu	Asp	Ser	Ser	Ala	Ala	Trp	Glu	Gln	Ser
65					70					75					80
Leu	Leu	Glu	Lys	Phe	Ser	Thr	Glu	Leu	Tyr	Gln	Gln	Leu	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Glu	Asp	Phe	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Met	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Lys	Lys
145					150					155					160
Gly	Leu	Arg	Arg	Lys	Asp										
				165											

<210> 37

<211> 166

<212> PRT

<213> Homo sapiens

<220>

<223> mature huIFN alpha-8b

<400> 37

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	Asn	Arg	Arg	Ala	Leu	Ile
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Glu	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Asp	Lys	Gln	Phe
		35					40					45			
Gln	Lys	Ala	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Leu	Asp	Glu	Thr
65					70					75					80
Leu	Leu	Asp	Glu	Phe	Tyr	Ile	Glu	Leu	Asp	Gln	Gln	Leu	Asn	Asp	Leu
				85					90					95	
Glu	Ser	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Ile	Glu	Ser	Pro	Leu	Met
			100					105					110		
Tyr	Glu	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Ser	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Ile	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Lys	Ser	Lys	Glu										
				165											

<210> 38

<211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-10a

<400> 38
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly Asn Arg Arg Ala Leu Ile
 1 5 10 15
 Leu Leu Gly Gln Met Gly Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Ile Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Glu Asp Ser Ser Ala Ala Trp Glu Gln Ser
 65 70 75 80
 Leu Leu Glu Lys Phe Ser Thr Glu Leu Tyr Gln Gln Leu Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Ile Glu Arg Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Leu Ser Phe Ser Thr Asn Leu Gln Lys
 145 150 155 160
 Arg Leu Arg Arg Lys Asp
 165

<210> 39
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-14a

<400> 39
 Cys Asn Leu Ser Gln Thr His Ser Leu Asn Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Glu Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asn Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Met Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys
 145 150 155 160

Arg Leu Arg Arg Lys Asp
165

<210> 40
<211> 166
<212> PRT
<213> Homo sapiens

<220>
<223> mature huIFN alpha-16

<400> 40
Cys Asp Leu Pro Gln Thr His Ser Leu Gly Asn Arg Arg Ala Leu Ile
1 5 10 15
Leu Leu Ala Gln Met Gly Arg Ile Ser His Phe Ser Cys Leu Lys Asp
20 25 30
Arg Tyr Asp Phe Gly Phe Pro Gln Glu Val Phe Asp Gly Asn Gln Phe
35 40 45
Gln Lys Ala Gln Ala Ile Ser Ala Phe His Glu Met Ile Gln Gln Thr
50 55 60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65 70 75 80
Leu Leu Asp Lys Phe Tyr Ile Glu Leu Phe Gln Gln Leu Asn Asp Leu
85 90 95
Glu Ala Cys Val Thr Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
100 105 110
Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
115 120 125
Leu Tyr Leu Met Gly Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
130 135 140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys
145 150 155 160
Gly Leu Arg Arg Lys Asp
165

<210> 41
<211> 166
<212> PRT
<213> Homo sapiens

<220>
<223> mature huIFN alpha-17b

<400> 41
Cys Asp Leu Pro Gln Thr His Ser Leu Gly Asn Arg Arg Ala Leu Ile
1 5 10 15
Leu Leu Ala Gln Met Gly Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
20 25 30
Arg His Asp Phe Gly Leu Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
35 40 45
Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln Thr
50 55 60
Phe Asn Leu Phe Ser Thr Glu Asp Ser Ser Ala Ala Trp Glu Gln Ser
65 70 75 80
Leu Leu Glu Lys Phe Ser Thr Glu Leu Tyr Gln Gln Leu Asn Asn Leu
85 90 95
Glu Ala Cys Val Ile Gln Glu Val Gly Met Glu Glu Thr Pro Leu Met
100 105 110
Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr

	115					120						125							
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val				
	130					135					140								
Arg	Ala	Glu	Ile	Met	Arg	Ser	Leu	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys				
145					150					155					160				
Ile	Leu	Arg	Arg	Lys	Asp														
				165															

<210> 42
 <211> 166
 <212> PRT
 <213> Homo sapiens

<220>
 <223> mature huIFN alpha-21

<400> 42																			
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	Asn	Arg	Arg	Ala	Leu	Ile				
1				5					10					15					
Leu	Leu	Ala	Gln	Met	Gly	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp				
		20						25					30						
Arg	His	Asp	Phe	Gly	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe				
	35					40						45							
Gln	Lys	Ala	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Ile	Gln	Gln	Thr				
	50					55					60								
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Thr	Trp	Glu	Gln	Ser				
65					70					75					80				
Leu	Leu	Glu	Lys	Phe	Ser	Thr	Glu	Leu	Asn	Gln	Gln	Leu	Asn	Asp	Leu				
			85						90					95					
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met				
		100						105					110						
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Lys	Lys	Tyr	Phe	Gln	Arg	Ile	Thr				
	115						120					125							
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val				
	130				135						140								
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Lys	Ile	Phe	Gln	Glu				
145				150						155					160				
Arg	Leu	Arg	Arg	Lys	Glu														
				165															

<210> 43
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha-Con1

<400> 43																			
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	Asn	Arg	Arg	Ala	Leu	Ile				
1				5					10					15					
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp				
		20						25					30						
Arg	His	Asp	Phe	Gly	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe				
	35					40						45							
Gln	Lys	Ala	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Ile	Gln	Gln	Thr				
	50					55					60								
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Ser				
65					70					75					80				

Leu	Leu	Glu	Lys	Phe	Tyr	Thr	Glu	Leu	Tyr	Gln	Gln	Leu	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Lys	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 44
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14C2a

<400> 44															
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25				30			
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40					45				
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55				60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asn	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65				70					75					80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
		100						105				110			
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115					120						125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 45
 <211> 167
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO1

<400> 45															
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25				30			
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55						60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
		100						105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130				135						140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 48
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CH05

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55						60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
		100						105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130				135						140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Cys	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 49
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>

<223> IFNalpha B9x14CHO6

<400> 49

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Cys	Lys	Glu										
				165											

<210> 50

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha 14Ep01

<400> 50

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	Gln	Phe
		35					40					45			
Gln	Lys	Ala	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Met	Met	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 51
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha 14Ep02

<400> 51
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Asp Lys Phe Tyr Ile Glu Leu Phe Gln Gln Leu Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Thr Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 52
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha 14Ep03

<400> 52
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140

Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 53
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha 14Ep04

<400> 53
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 54
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha 14Ep05

<400> 54
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn Gln Phe
 35 40 45
 Gln Lys Ala Gln Ala Ile Ser Val Leu His Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met

Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 57

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x14CH04C31

<400> 57

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Cys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55				60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 58

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x14CH04C46

<400> 58

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp

<400> 60

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55						60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Cys	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130				135						140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 61

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x14CH04C79

<400> 61

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55						60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Cys	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
	115						120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130				135						140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 62

<211> 166

<212> PRT
<213> Artificial Sequence

<220>
<223> IFNalpha B9x14CH04C107

<400> 62
Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
1 5 10 15
Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
20 25 30
Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
35 40 45
Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
50 55 60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65 70 75 80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
85 90 95
Glu Ala Cys Val Met Gln Glu Val Gly Val Cys Glu Thr Pro Leu Met
100 105 110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
115 120 125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
130 135 140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
145 150 155 160
Ser Leu Arg Ser Lys Glu
165

<210> 63
<211> 166
<212> PRT
<213> Artificial Sequence

<220>
<223> IFNalpha B9x14CH04C122

<400> 63
Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
1 5 10 15
Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
20 25 30
Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
35 40 45
Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
50 55 60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65 70 75 80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
85 90 95
Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
100 105 110
Asn Val Asp Ser Ile Leu Ala Val Arg Cys Tyr Phe Gln Arg Ile Thr
115 120 125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
130 135 140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
145 150 155 160
Ser Leu Arg Ser Lys Glu

<210> 64
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CH04C134

<400> 64
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Cys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 65
 <211> 160
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14Ep04(161-166)

<400> 65
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125

Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160

<210> 66
 <211> 164
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14Ep04(165-166

<400> 66
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser

<210> 67
 <211> 159
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14Ep04(1-4D44*(161-166

<400> 67
 Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met Leu Leu Ala Gln
 1 5 10 15
 Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp Arg His Asp Phe
 20 25 30
 Arg Phe Pro Gln Glu Glu Phe Gly Asn His Phe Gln Lys Val Gln Ala
 35 40 45
 Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr Phe Asn Leu Phe Ser
 50 55 60
 Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr Leu Leu Glu Lys Phe
 65 70 75 80
 Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu Glu Ala Cys Val Met
 85 90 95
 Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met Asn Val Asp Ser Ile

Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 70
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO8

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Arg	Asp
		20						25				30			
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
	35					40					45				
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50				55					60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65				70					75					80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85					90						95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105				110			
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 71
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO9

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25				30			
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Arg	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
				85					90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 74
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO12

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
		20						25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
	50					55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Arg	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 75
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>

<223> IFNalpha B9x14CHO13

<400> 75

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Arg	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 76

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x14CHO14

<400> 76

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Gly	His	Arg	Arg	Thr	Met	Met
1				5					10					15	
Leu	Leu	Ala	Gln	Met	Arg	Arg	Ile	Ser	Leu	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Arg	Phe	Pro	Gln	Glu	Glu	Phe	Asp	Gly	Asn	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Phe	Tyr	Glu	Met	Met	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Arg	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 77
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO15

<400> 77
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Arg Glu
 165

<210> 78
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CHO16

<400> 78
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Arg Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Arg Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140

Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 79
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CH017

<400> 79
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Arg Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Arg Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 80
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x14CH018

<400> 80
 Cys Asp Leu Pro Gln Thr His Ser Leu Gly His Arg Arg Thr Met Met
 1 5 10 15
 Leu Leu Ala Gln Met Arg Arg Ile Ser Leu Phe Ser Cys Leu Arg Asp
 20 25 30
 Arg His Asp Phe Arg Phe Pro Gln Glu Glu Phe Asp Gly Asn His Phe
 35 40 45
 Gln Lys Val Gln Ala Ile Phe Leu Phe Tyr Glu Met Met Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Met Gln Glu Val Gly Val Glu Glu Thr Pro Leu Met

Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asp	Leu
			85						90					95	
Glu	Ala	Cys	Val	Met	Gln	Glu	Val	Gly	Val	Glu	Glu	Thr	Pro	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Arg	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser												

<210> 83
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25CHO1

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35				40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
	50				55					60					
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
			85						90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
	130					135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Phe	Ser	Thr	Asn	Leu	Gln	Lys
145					150					155					160
Arg	Leu	Arg	Arg	Lys	Glu										
				165											

<210> 84
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25CHO2

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp

<400> 86

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Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1          5          10          15
Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
          20          25          30
Arg His Asp Phe Gly Phe Pro Glu Glu Phe Asp Gly His His Phe
          35          40          45
Gln Lys Val Gln Ala Ile Phe Leu Leu Tyr Glu Leu Ile Gln Gln Thr
          50          55          60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65          70          75          80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
          85          90          95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
          100          105          110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
          115          120          125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
          130          135          140
Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
145          150          155          160
Ser Leu Arg Cys Lys Glu
          165
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<210> 87

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x25Ep01

<400> 87

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Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1          5          10          15
Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
          20          25          30
Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
          35          40          45
Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
          50          55          60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65          70          75          80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
          85          90          95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
          100          105          110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
          115          120          125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
          130          135          140
Arg Ala Glu Ile Met Arg Ser Phe Ser Phe Ser Thr Asn Leu Gln Lys
145          150          155          160
Arg Leu Arg Arg Lys Glu
          165
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<210> 88

<211> 166

<212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep02

<400> 88
 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Ile Met Ala Gln Met Gly Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 89
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep03

<400> 89
 Cys Asn Leu Ser Gln Thr His Ser Leu Asn Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu

<210> 90
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep04

<400> 90
 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Met Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 91
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep05

<400> 91
 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125

Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 92
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep06

<400> 92
 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Asp Lys Phe Tyr Ile Glu Leu Phe Gln Gln Leu Asn Asp Leu
 85 90 95
 Glu Ala Cys Val Thr Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Arg Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165

<210> 93
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep07

<400> 93
 Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
 1 5 10 15
 Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
 20 25 30
 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu

				85					90				95			
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met	
			100					105					110			
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr	
		115					120					125				
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val	
	130				135						140					
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu	
145				150						155					160	
Ser	Leu	Arg	Ser	Lys	Glu											
				165												

<210> 94
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep08

<400> 94																
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met	
1				5					10					15		
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp	
		20						25				30				
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe	
	35					40					45					
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr	
	50				55					60						
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr	
65				70					75					80		
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu	
			85					90					95			
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met	
		100						105				110				
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr	
	115					120						125				
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val	
	130				135					140						
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu	
145				150						155					160	
Ser	Leu	Arg	Ser	Lys	Glu											
				165												

<210> 95
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep10

<400> 95																
Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met	
1				5					10					15		
Ile	Met	Ala	Gln	Met	Gly	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp	
		20						25				30				
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe	
	35					40					45					

Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
50						55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
			85						90				95		
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
			115				120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
			130			135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 96

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x25Ep11

<400> 96

Cys	Asn	Leu	Ser	Gln	Thr	His	Ser	Leu	Asn	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35				40						45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
			85						90				95		
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
			115				120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
			130			135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 97

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x25Ep12

<400> 97

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

1	5	10	15
Leu Met Ala Gln Met Arg Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp			
	20	25	30
Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His His Phe			
	35	40	45
Gln Lys Val Gln Ala Ile Phe Leu Leu Tyr Glu Leu Ile Gln Gln Thr			
	50	55	60
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr			
65	70	75	80
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu			
	85	90	95
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met			
	100	105	110
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr			
	115	120	125
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val			
	130	135	140
Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu			
145	150	155	160
Ser Leu Arg Ser Lys Glu			
	165		

<210> 98
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> IFNalpha B9x25Ep13

<400> 98
Cys Asp Leu Pro Gln Thr His Ser Leu Ser Asn Arg Arg Thr Leu Met
1
Ile Met Ala Gln Met Gly Arg Ile Ser Pro Phe Ser Cys Leu Lys Asp
20
Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His His Phe
35
Gln Lys Val Gln Ala Ile Phe Leu Leu Tyr Glu Leu Ile Gln Gln Thr
50
Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
65
Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
85
Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
100
Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
115
Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
130
Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
145
150
Ser Leu Arg Ser Lys Glu
165

<210> 99
 <211> 166
 <212> PRT
 <213> Artificial Sequence

<220>

<223> IFNalpha B9x25Ep14

<400> 99

Cys	Asn	Leu	Ser	Gln	Thr	His	Ser	Leu	Asn	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Glu	Phe	Asp	Gly	His	His	Phe
		35					40					45			
Gln	Lys	Val	Gln	Ala	Ile	Phe	Leu	Leu	Tyr	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
			85						90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Gln	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
145					150					155					160
Ser	Leu	Arg	Ser	Lys	Glu										
				165											

<210> 100

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> IFNalpha B9x25Ep15

<400> 100

Cys	Asp	Leu	Pro	Gln	Thr	His	Ser	Leu	Ser	Asn	Arg	Arg	Thr	Leu	Met
1				5					10					15	
Leu	Met	Ala	Gln	Met	Arg	Arg	Ile	Ser	Pro	Phe	Ser	Cys	Leu	Lys	Asp
			20					25					30		
Arg	His	Asp	Phe	Gly	Phe	Pro	Glu	Glu	Phe	Asp	Gly	His	Gln	Phe	
		35					40				45				
Gln	Lys	Thr	Gln	Ala	Ile	Ser	Val	Leu	His	Glu	Leu	Ile	Gln	Gln	Thr
		50				55					60				
Phe	Asn	Leu	Phe	Ser	Thr	Lys	Asp	Ser	Ser	Ala	Ala	Trp	Asp	Glu	Thr
65					70					75				80	
Leu	Leu	Glu	Lys	Phe	Tyr	Ile	Glu	Leu	Phe	Gln	Gln	Met	Asn	Asn	Leu
			85						90					95	
Glu	Ala	Cys	Val	Ile	Gln	Glu	Val	Gly	Val	Glu	Glu	Ile	Ala	Leu	Met
			100					105					110		
Asn	Val	Asp	Ser	Ile	Leu	Ala	Val	Arg	Lys	Tyr	Phe	Arg	Arg	Ile	Thr
		115					120					125			
Leu	Tyr	Leu	Thr	Glu	Lys	Lys	Tyr	Ser	Pro	Cys	Ala	Trp	Glu	Val	Val
		130				135					140				
Arg	Ala	Glu	Ile	Met	Arg	Ser	Phe	Ser	Leu	Ser	Thr	Asn	Leu	Gln	Glu
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 Arg His Asp Phe Gly Phe Pro Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
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 Ser Leu Arg Ser Lys Glu
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 Arg His Asp Phe Gly Phe Pro Glu Glu Glu Phe Asp Gly His Gln Phe
 35 40 45
 Gln Lys Thr Gln Ala Ile Ser Val Leu His Glu Leu Ile Gln Gln Thr
 50 55 60
 Phe Asn Leu Phe Ser Thr Lys Asp Ser Ser Ala Ala Trp Asp Glu Thr
 65 70 75 80
 Leu Leu Glu Lys Phe Tyr Ile Glu Leu Phe Gln Gln Met Asn Asn Leu
 85 90 95
 Glu Ala Cys Val Ile Gln Glu Val Gly Val Glu Glu Ile Ala Leu Met
 100 105 110
 Asn Val Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu

Asn Glu Asp Ser Ile Leu Ala Val Arg Lys Tyr Phe Gln Arg Ile Thr
 115 120 125
 Leu Tyr Leu Thr Glu Lys Lys Tyr Ser Pro Cys Ala Trp Glu Val Val
 130 135 140
 Arg Ala Glu Ile Met Arg Ser Phe Ser Leu Ser Thr Asn Leu Gln Glu
 145 150 155 160
 Ser Leu Arg Ser Lys Glu
 165